## Introduction to Data Science

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## **DISCUSSION**

What do we mean by data having structure?

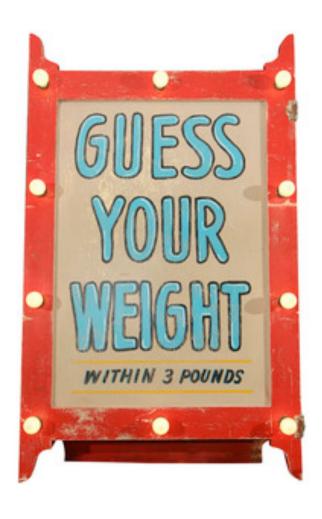


Why do we want to understand it?

What do we do with it?

### INFORMATION

# any quantity that can reduce uncertainty about another quantity

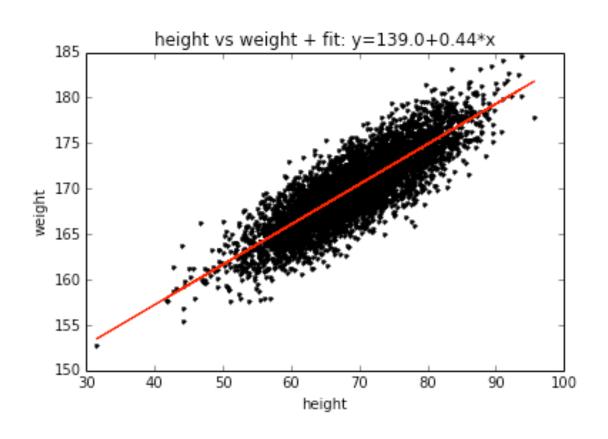


If a carnival operator wanted to reduce the uncertainty about your weight, what information might he use?

## REDUCING UNCERTAINTY

#### Let's assume our carnival friend is also a data scientist

Step 1 – Collect data, height vs. weight, regress



By regressing we can learn:

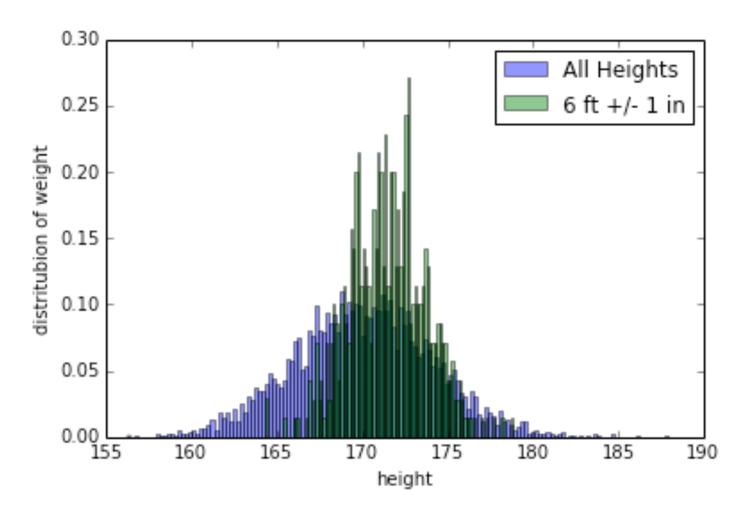
E[Weight|Height]

as opposed to

E[Weight].

## REDUCING UNCERTAINTY

Step 2 – Based on prior knowledge, use the person's height to get a better range of possible weights.

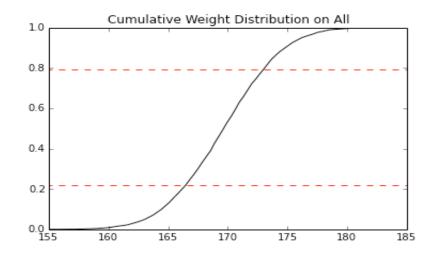


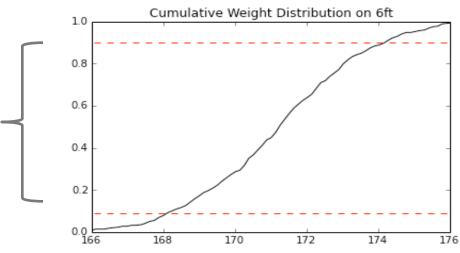
## REDUCING UNCERTAINTY

Step 3 – Guess E[W|H=72]. With a more informed, conditional expectation, he can reduce losses by 50% (40/100 vs. 20/100). That could be a lot of \$\$\$.



*Error Rate* = *P(Correct* | *Guess E[W]*|*H*=72) = 20%





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# EXPLORATORY ANALYSIS: FINDING STRUCTURE

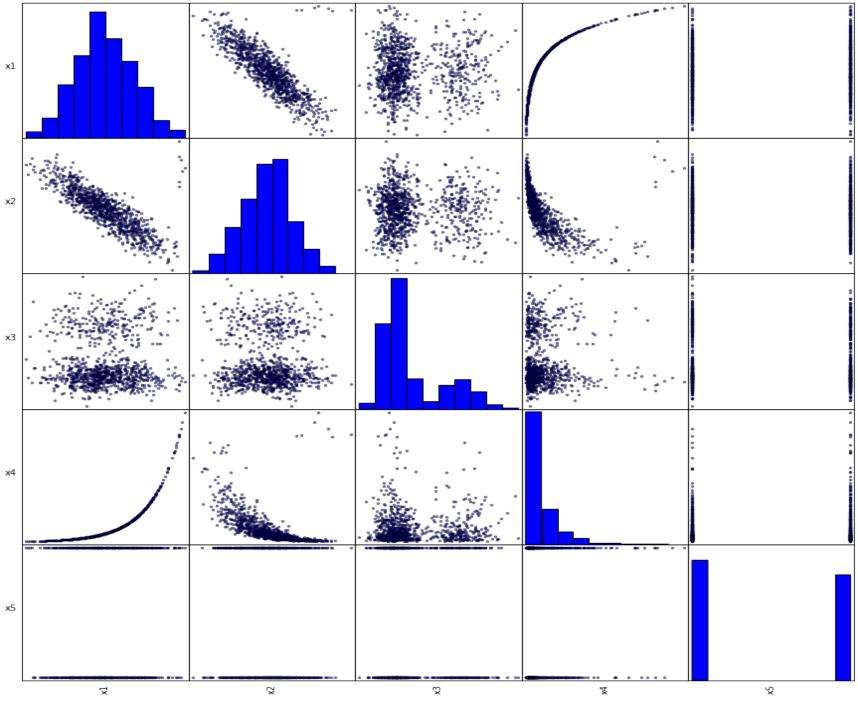
## **TYPES OF STRUCTURE**

<u>Univariate</u> – does a single variable follow a predictable pattern?

Pairwise (i.e., between 2 variables)

Global (i.e., across a matrix, or within a set of variables)

**Supervised** (technically includes the above, but with special emphasis on a single target variable)



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## **TYPES OF STRUCTURE**

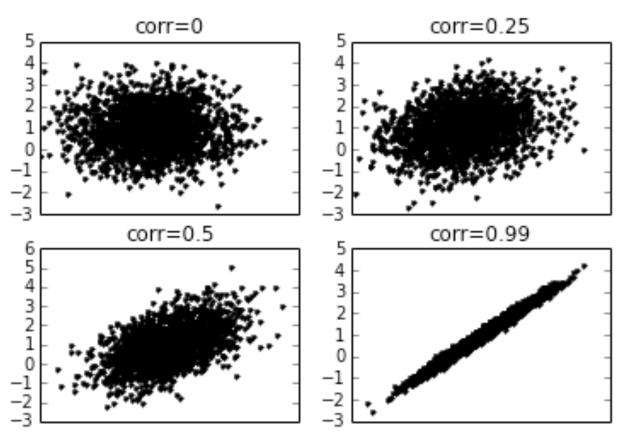


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### **COVARIANCE AND CORRELATION**

These are probably the most used and thought of metrics when considering pairwise structure. These are statistical quantities and have a fairly intuitive geometric interpretation.

#### Scatter of Bi-Variate Normally Distributed Variables with various Correlations



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## COVARIANCE

#### Covariance

$$\begin{split} \sigma(x,y) &= \mathrm{E}\left[\left(x-\mathrm{E}\left[x\right]\right)\left(y-\mathrm{E}\left[y\right]\right)\right] \\ &= \mathrm{E}\left[xy-x\,\mathrm{E}\left[y\right]-\mathrm{E}\left[x\right]y+\mathrm{E}\left[x\right]\mathrm{E}\left[y\right]\right] \\ &= \mathrm{E}\left[xy\right]-\mathrm{E}\left[x\right]\mathrm{E}\left[y\right]-\mathrm{E}\left[x\right]\mathrm{E}\left[y\right]+\mathrm{E}\left[x\right]\mathrm{E}\left[y\right] \\ &= \mathrm{E}\left[xy\right]-\mathrm{E}\left[x\right]\mathrm{E}\left[y\right]. \end{split}$$

#### Sample Covariance (between Xj and Xk)

$$q_{jk} = \frac{1}{N-1} \sum_{i=1}^{N} (x_{ij} - \bar{x}_j) (x_{ik} - \bar{x}_k)$$

Note: I have decided to pull equations often from wikipedia, as I feel wp represents a crowd-sourced vote on notation standardization: http://en.wikipedia.org/wiki/Covariance

## CORRELATION

Aka: Pearson-Product Moment Correlation Coefficient. This is just the covariance normalized by the variance of each variable. This scales correlation to the interval [-1,1], which makes it a very intuitive tool for analysis and reporting.

$$\rho_{X,Y} = \operatorname{corr}(X,Y) = \frac{\operatorname{cov}(X,Y)}{\sigma_X \sigma_Y} = \frac{E[(X - \mu_X)(Y - \mu_Y)]}{\sigma_X \sigma_Y},$$

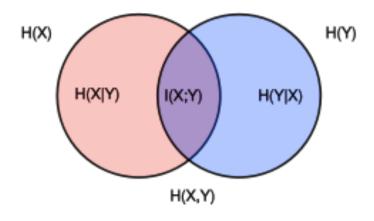
**Sample Correlation** 

$$r_{xy} = \frac{\sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})}{(n-1)s_x s_y} = \frac{\sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^{n} (x_i - \bar{x})^2 \sum_{i=1}^{n} (y_i - \bar{y})^2}},$$

Almost any programming language has standard functions for these formulas, but it doesn't hurt to understand them!

Equation Source: http://en.wikipedia.org/wiki/Covariance

This comes from Information Theory, is used often for feature importance ranking and is related to important aspects of Decision Tree algorithms.



#### **Mutual Information**

$$I(X;Y) = \sum_{y \in Y} \sum_{x \in X} p(x,y) \log \left( \frac{p(x,y)}{p(x) p(y)} \right),$$

Let's break this down

#### **Mutual Information**

$$I(X;Y) = \sum_{y \in Y} \sum_{x \in X} p(x,y) \log \left( \frac{p(x,y)}{p(x) p(y)} \right),$$

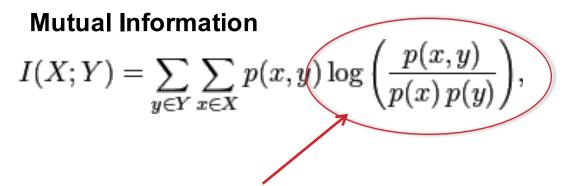
This is in the form of E[F], i.e., E[X]= $\Sigma p(F)^*F$ . In this case, what is X?

Let's break this down

Mutual Information  $I(X;Y) = \sum_{y \in Y} \sum_{x \in X} p(x,y) \log \left(\frac{p(x,y)}{p(x) \ p(y)}\right),$ 

F=log(p(x,y)/(p(x)\*p(y))....but what is this?

Let's break this down



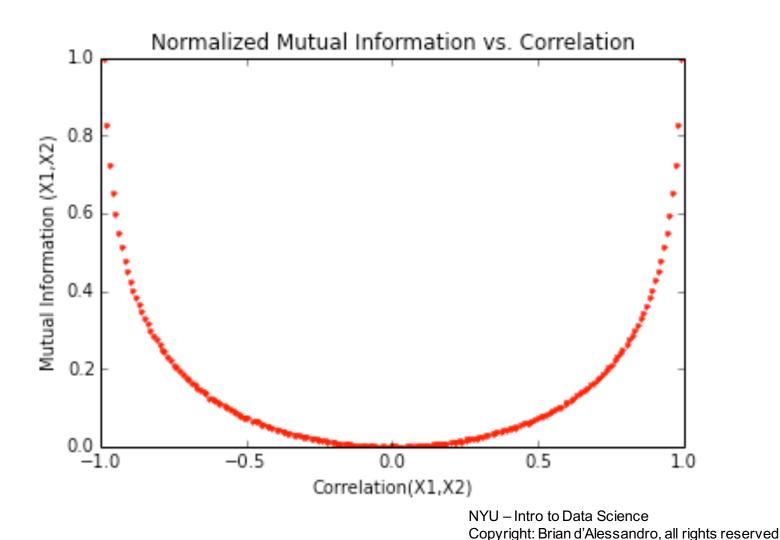
F=log(p(x,y)/(p(x)\*p(y))....but what is this?

This is a quantity with the following properties:

- If X,Y are independent, F=0
- If X,Y are completely dependent, F is at a maximum
- F is symmetric (F(x,y)=F(y,x))

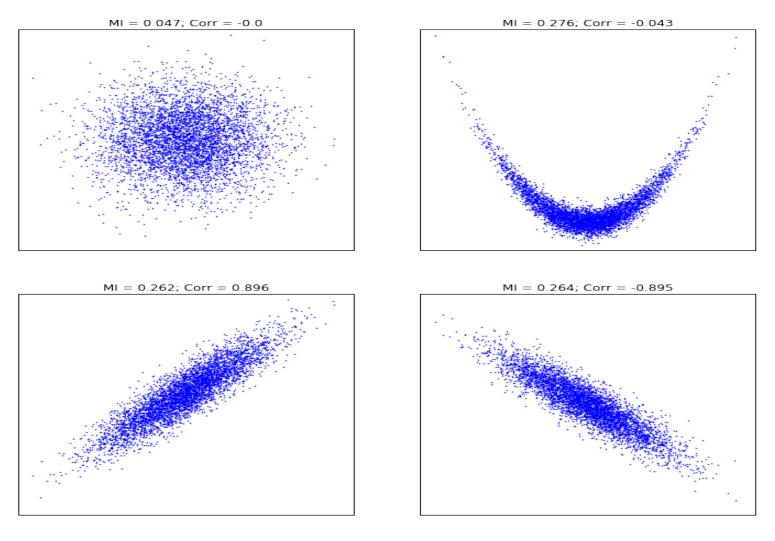
## MI VS CORRELATION

Scikit-learn has functions for MI and normalized Mutual Information. We can see that MI and correlation are monotonically related concepts, though MI is strictly positive so does not indicate negative dependencies.



## MI VS CORRELATION

We can see the difference by looking at both quantities for different types of variable dependencies.



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## MI VS CORRELATION

#### **Mutual Information**

- Can capture non-linear dependencies better
- Works naturally with categorical data

#### **Correlation Coefficient**

- Expresses negative dependencies
- Well understood and intuitive (easy to communicate)

### **PUTTING THESE TO USE**

- General understanding of dependencies in data
- Validating assumptions for statistical modeling
- Feature ranking and selection
- Decision Tree Algorithms

## **TYPES OF STRUCTURE**



**Global** (i.e., across a matrix, or within a set of variables)

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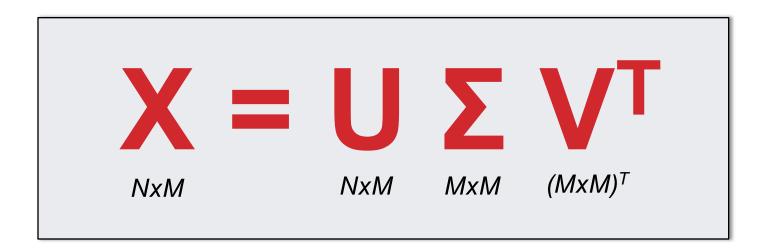
## DATA IS MULTIVARIATE

We usually want to go beyond pairwise similarity and understand how much "information" is actually embedded in a matrix.

We also might want to know what groups of features are related.

In an Nx2 matrix, this problem reduces to the pairwise methods just discussed.

Although this is not a linear algebra course, this equation happens so often in data analysis that it is worth learning (again).



We will not dive into all of the theoretical aspects of SVD and related topics. Our goal here is to understand it intuitively and be able to use it as a tool for solving other common Data Science problems.

Lets go through this piece by piece:

U

U holds the left singular vectors. Each row contains k elements that correspond to the latent factors of each row of X. U is orthonormal.

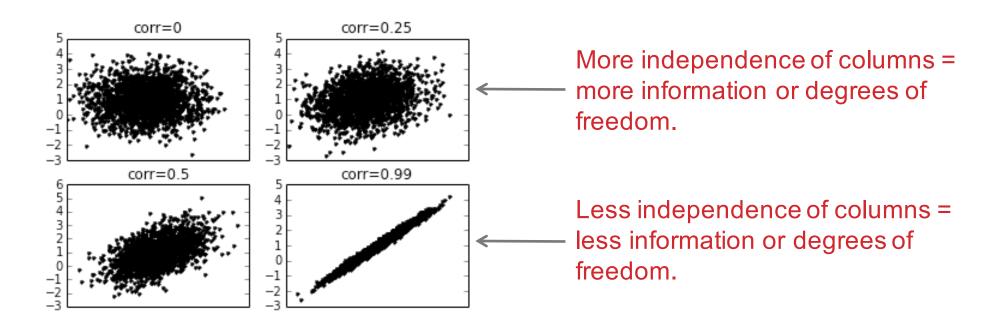
 $\Sigma$  is a diagonal matrix that holds the singular values (in descending order). The singular values are essentially weights that determine how much that latent factor contributes to the matrix.



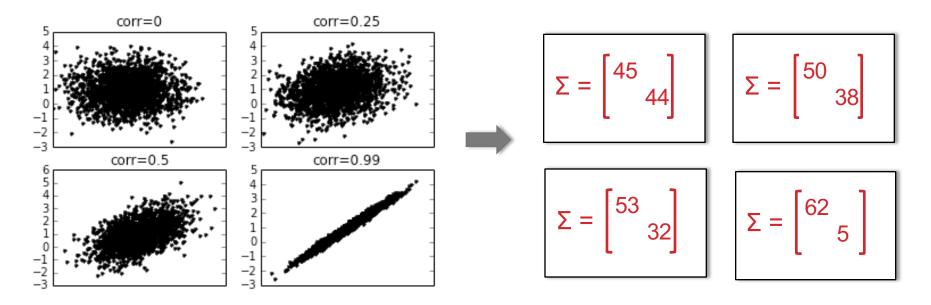
V holds the right singular vectors. Each row contains k elements that correspond to the latent factors of each column of X. V is orthonormal.

How does this relate to structure?

Lets recall these scatter plots. Each plot can be expressed as an Nx2 Matrix. The SVD is a tool that can help us understand how much information or structure is actually in the matrix.



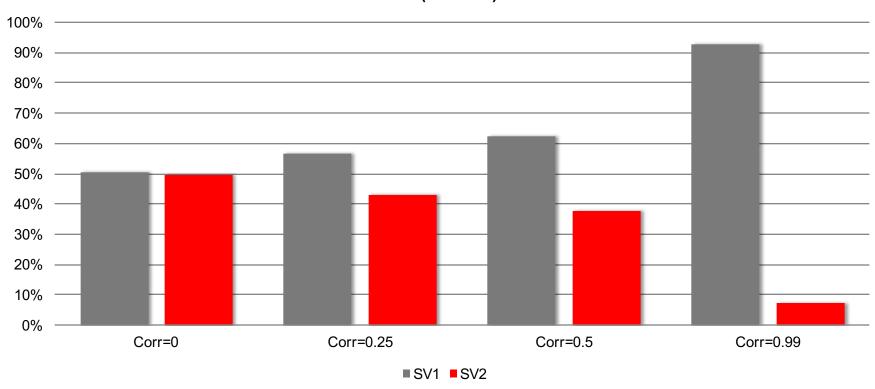
Using Scipy we can easily compute the SVD.: U,Sig,Vt = scipy.linalg.svd(matrix)



The magnitude of the singular-values are generally determined by the magnitude of the values in X. The relative difference between singular values is a function of the level of independence of the columns.

We can see that less independence of the columns leads to singular values with more skew from each other.

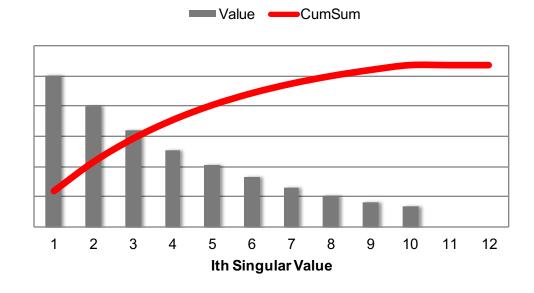




This idea generalizes to more dimensions, which is where the SVD is extremely useful.

The skew of the singular values, and the shape of the cumulative sum curve can give us a sense of the degree of independence in a multi-dimensional matrix.

#### **Summary of Singular Values**



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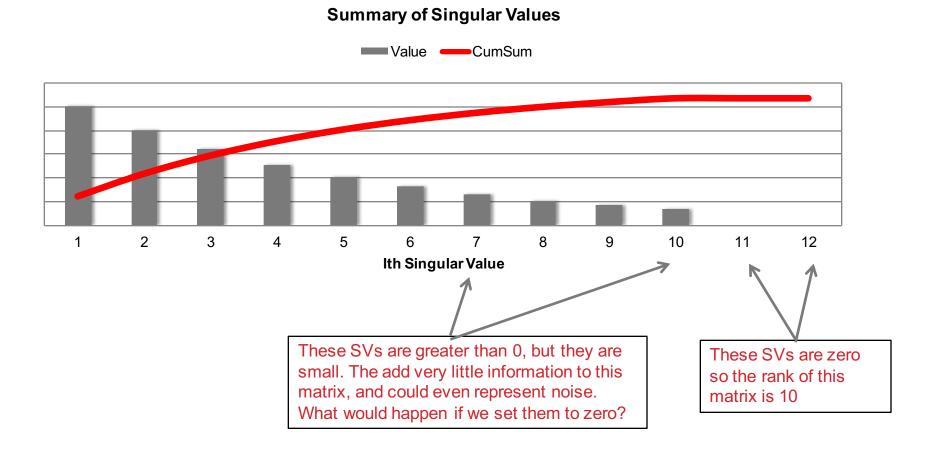
# AN INCREDIBLY USEFUL APPLICATION OF SVD

One of the most powerful applications of the SVD is creating a low rank approximation of a data matrix. Many of the following methods are based on using the SVD to get a low rank approximation.

- Data Compression
- Dimensionality Reduction
- Recommender Systems
- Clustering in High Dimensions

### THE LOW RANK APPROXIMATION

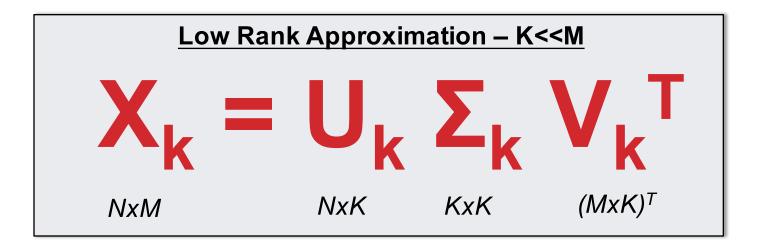
The rank of a matrix is the size of the largest number of independent columns of a matrix. The rank can be found by counting the number of singular values >0.



#### THE LOW RANK APPROXIMATION

We can build a matrix  $X_k$  that approximates our original matrix by doing the following:

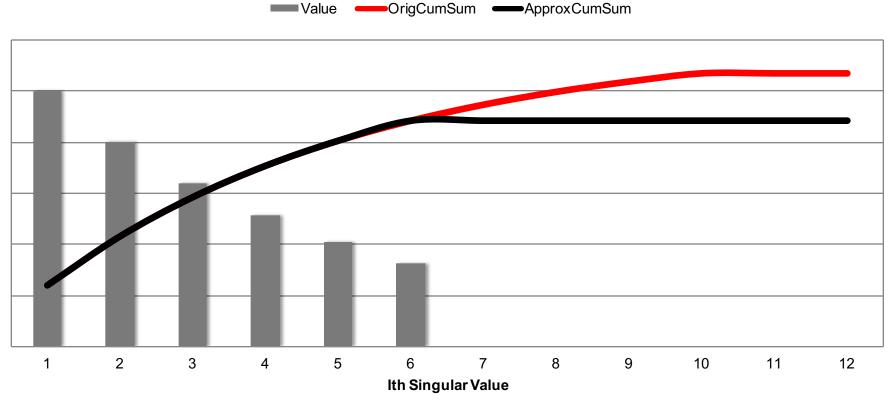
- 1. Compute the SVD of X
- 2. Truncate U, Σ and V to take only the k highest columns & singular values
- 3. Multiply back together to get  $X_k$



#### THE LOW RANK APPROXIMATION

Our original matrix had 12 columns with a rank of 10. In our approximation, we decided to use k=6. The cumulative sum of singular values is related to the amount of information contained in the matrix. By using k=6, we lost some information, but not half.

## What do we gain with the low rank approximation? What do we lose?



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## ONE BENEFIT OF RANK-K SVD

#### **Some facts**

- A floating point number uses 8 bytes of memory
- An MxN matrix of floating point numbers uses 8\*M\*N bytes of memory.

Instead of storing X, lets use  $X_k$ , a low rank approximation

 $oldsymbol{\mathsf{U}_k}$  ...this is 8\*N\*K bytes When separated...  $oldsymbol{\mathsf{\Sigma}_k}$  ...this is 8\*1\*K bytes  $oldsymbol{\mathsf{8*K*(N+M+1)}}$  bytes  $oldsymbol{\mathsf{V_k^T}}$  ...this is 8\*K\*K bytes

If you choose K<<M, you could save a ton of space...

i.e. k=100,N=1MM,M=10k, the full X is 80 GB, whereas storing the decomposed components of  $X_k$  would only be 0.8 GB!

## THE COST

This result comes from the Eckart-Young-Mirsky Theorem

If X is an nxm matrix and  $X_k$  is the rank-k approximation derived from the SVD, then the sum-of-squares error between the entries of X and  $X_k$  equals the square-root of the sum-of-squares of the singular values > k.

i.e.

$$||X - X_k||_F = \sqrt{\sum_{r=k+1}^{\min(m,n)} \sigma_r^2}$$

Where the Frobenius norm is defined as:

$$||A||_F = \sqrt{\sum_{i=1}^n \sum_{j=1}^m a_{ij}^2}$$

### THE COST - EXAMPLE

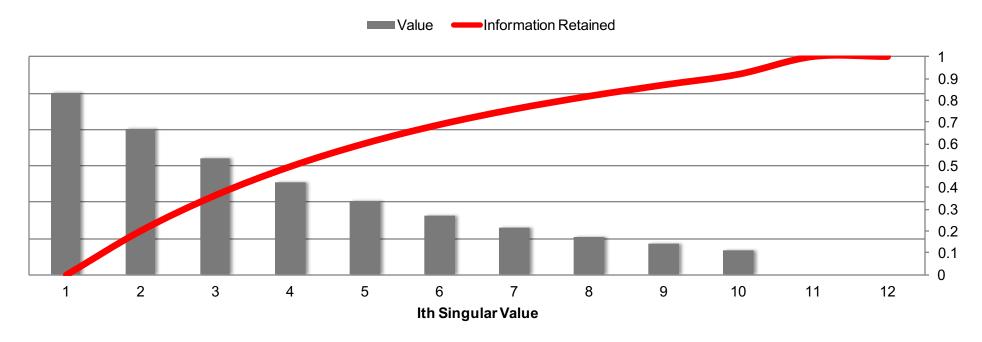
Given a matrix with the Singular Values shown below, the "Information Retained" shows:

$$1 - \frac{\|X - X_k\|_F}{\|X\|_F}$$

This is very similar to the  $R^2$  metric in linear regression, and it tells us how well our rank-k approximation "fits" the the original matrix from a least squares sense.

Analyzing a rank-*k* approximation like this gives us a tool to choose *k*.

#### **Summary of Singular Values**



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## **DIMENSIONALITY REDUCTION**

Sometimes a dataset has too many features for various algorithms (or storage systems to handle). Often times there is also a lot of redundancy of the data. The SVD gives us a way to reduce the number of variables without losing too much information.

Compute rank-k SVD 
$$X_k = U_k \Sigma_k V_k^T$$

We can create an NxK reduced matrix with  $XV_k$  This effectively becomes our new "X" matrix, and we do any analysis (clustering, modeling, etc.) with the reduced matrix.

We can project any new data into the reduced space by,  $\mathbf{X}' \mathbf{V}_{\mathbf{k}}$  and then use this in our algorithms.

Again, the optimal choice of K is dependent on the problem and will be a tradeoff between information loss vs. constraint tolerance.

# IMAGE COMPRESSION EXAMPLE IN PYTHON

Images can be thought of as matrices of pixel color values. We can use SVD to build faces with different ranks. The full matrix here has rank=200 but we can see with a rank as low as 10 we capture most of the information we need to recognize the image.



## **BUILDING AN EFFICIENT LOOKUP TABLE – THE CONTEXT**

At Dstillery, we operate a mobile advertising bidding system.

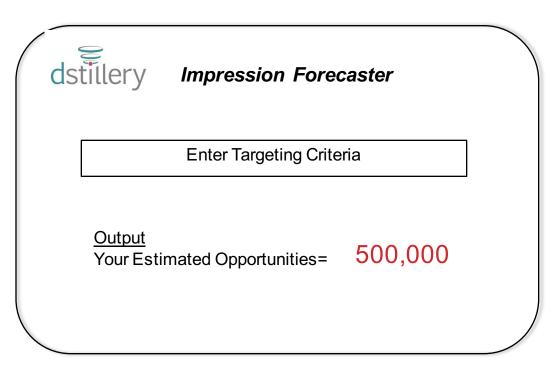
A user initiates an ad call by using an App. The Exchange receives the ad call and passes it to Dstillery as an option to bid.



If Dstillery places the highest bid, it sends an ad unit to the exchange to be displayed on the phone.

# **BUILDING AN EFFICIENT LOOKUP TABLE – THE PROBLEM**

Dstillery would like to provide its clients with a web application for forecasting total impression opportunities in a given period. The client would enter a set of targeting criteria (location, app name, device type, etc.) and Dstillery would provide an estimate.



Seems simple enough, but wait!

# **BUILDING AN EFFICIENT LOOKUP TABLE – THE NAÏVE SOLUTION**

An easy, brute force way to forecast is to just count the number of bid opportunities that happened in the past and use that as a forecast for the future.

State	Zip	Census	Арр	Device Type	Cnt
NY	11222	1001	Candy	Phone	678
NY	10003	1002	HideMe	Phone	77,794
NJ	08096	2001	Whoa	Tablet	78,143
NJ	08910	2002	Stoopid	Phone	1,112
PA	07919	3001	Flava	Tablet	59,787
PA	09199	3002	BigStuff	Tablet	5,000
TX	77450	9982	Scream	Phone	66,953

This is easy to accomplish in a single map-reduce query. The challenge is we end up with a table that has ~ 100 MM rows. The problem is that:

Querying a 100 MM record table with a live web interface is way too slow!

## BUILDING AN EFFICIENT LOOKUP TABLE – EFFICIENT SOLUTION (1)

Step 1: Convert our table from a compound key-value format to a matrix.

State	Zip	Census	Арр	Device Type	Cnt
NY	11222	1001	Candy	Phone	678
NY	10003	1002	HideMe	Phone	77,794
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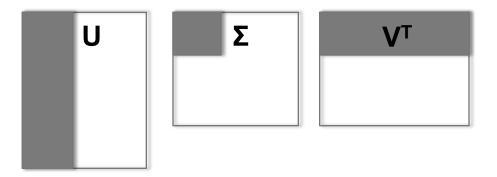




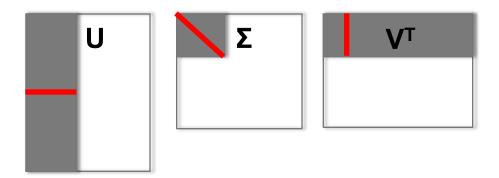
	App-Device 1	App-Device 2	 App-Device M
State,Zip,Census 1	1000	10,000	 90
State,Zip,Census 2	500	4000	 900
State,Zip,Census 3	2400	34000	 800
State,Zip,Census 4	900	10,000	 50
State,Zip,Census N	50	900	 100

## BUILDING AN EFFICIENT LOOKUP TABLE - EFFICIENT SOLUTION (2)

Step 2: Use SVD to decompose the matrix...store the top k rows/columns of the decomposed matrices.

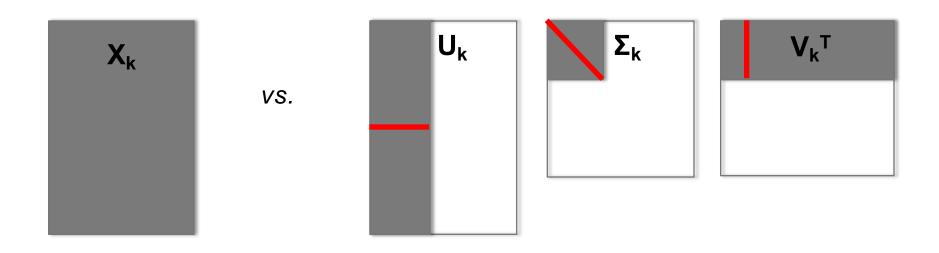


Step 3: During query time, look up appropriate row/columns from decomposed matrix. Multiply to get estimate:  $Forecast = \sum \sigma_i u_i v_i$ 



## **BUILDING AN EFFICIENT LOOKUP TABLE – COST/BENEFIT ANALYSIS**

- 1. The original key-value table had 100MM rows
- 2. For U and V we have K, 10k element vectors
- 3. Each scan of original table costs O(100MM)
- 4. Scanning U and V costs 2\*K\*O(10k)
- 5. Business requirement on accuracy is that we are right in the order of magnitude (i.e., log-scale), which means we can keep K<<10k



### TO BE CONTINUED...

Other applications of SVD are in recommender systems as well as clustering.

These will be presented in a later lecture.